

The Association of Hippocampal Long-Term Potentiation-Induced Gene Expression with Genetic Risk for Psychosis

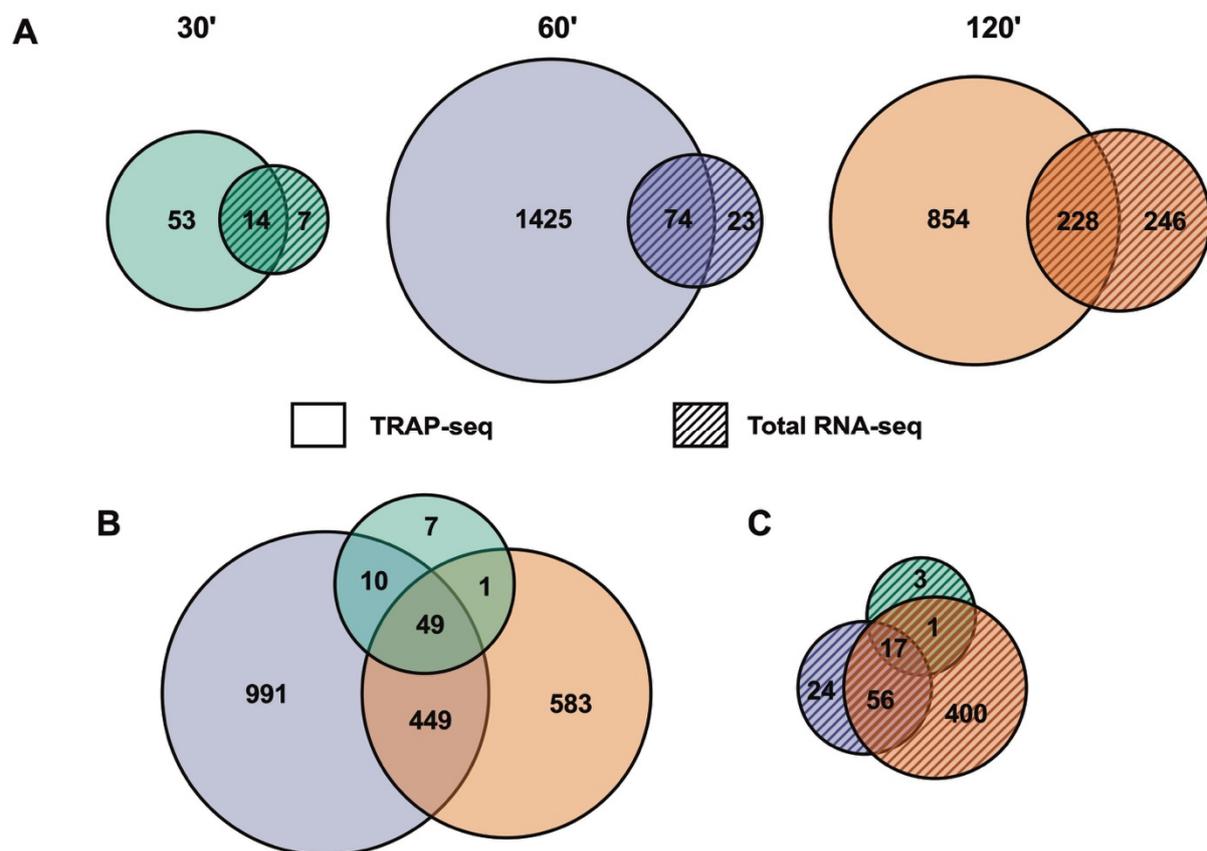
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Supplementary Information



Supplementary Figure S1 The intersection between genes differentially expressed (FDR < 0.01) in the TRAP and total sequencing gene sets. A. The number of genes differentially expressed at 30 min (green), 60 min (purple) and 120 min (orange). B. The overlap between genes differentially expressed in the TRAP-seq condition across time points. C. The overlap between genes differentially expressed in the total RNA-seq condition across time points. Solid fill represents the TRAP-seq condition, patterned fill represents the total RNA-seq condition. Not to scale.